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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/996,015

DATE: 05/20/2002

TIME: 15:54:52

Input Set : A:\Cg-81cil.app

Output Set: N:\CRF3\05202002\I996015.raw

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3 <110> APPLICANT: Quinn, Kerry E.
4   Pena, Carol A. E.
5   Li, Li
6   Spaderna, Steven K.
7   Leite, Marlo W.
9 <120> TITLE OF INVENTION: Aortic Carboxypeptidase-Like Proteins and Nucleic Acids
10  Encoding Same
12 <130> FILE REFERENCE: 15966-581 CIP
14 <140> CURRENT APPLICATION NUMBER: 09/996,015
15 <141> CURRENT FILING DATE: 2001-11-28
17 <150> PRIOR APPLICATION NUMBER: 09/641,741
18 <151> PRIOR FILING DATE: 2000-08-18
20 <150> PRIOR APPLICATION NUMBER: 60/159,613
21 <151> PRIOR FILING DATE: 1999-10-14
23 <150> PRIOR APPLICATION NUMBER: 60/175,534
24 <151> PRIOR FILING DATE: 2000-01-11
26 <150> PRIOR APPLICATION NUMBER: 60/224,086
27 <151> PRIOR FILING DATE: 2000-08-09
29 <160> NUMBER OF SEQ ID NOS: 47
31 <170> SOFTWARE: PatentIn Ver. 2.1
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34 <211> LENGTH: 2382
35 <212> TYPE: DNA
36 <213> ORGANISM: Homo sapiens
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81 <210> SEQ ID NO: 2

82 <211> LENGTH: 734

83 <212> TYPE: PRT

84 <213> ORGANISM: Homo sapiens

86 <400> SEQUENCE: 2

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91 20 25 30
93 Gly Thr Thr Lys Val Pro Gly Ser Thr Pro Ala Leu His Ser Ser Pro
94 35 40 45
96 Ala Gln Pro Pro Ala Glu Thr Ala Asn Gly Thr Ser Glu Gln His Val
97 50 55 60
99 Arg Ile Arg Val Ile Lys Lys Lys Val Ile Met Lys Lys Arg Lys
100 65 70 75 80
102 Lys Leu Thr Leu Thr Arg Pro Thr Pro Leu Val Thr Ala Gly Pro Leu
103 85 90 95
105 Val Thr Pro Thr Pro Ala Gly Thr Leu Asp Pro Ala Glu Lys Gln Glu
106 100 105 110
108 Thr Gly Cys Pro Pro Leu Gly Leu Glu Ser Leu Arg Val Ser Asp Ser
109 115 120 125
111 Arg Leu Glu Ala Ser Ser Ser Gln Ser Phe Gly Leu Gly Pro His Arg
112 130 135 140
114 Gly Arg Leu Asn Ile Gln Ser Gly Leu Glu Asp Gly Asp Leu Tyr Asp
115 145 150 155 160
117 Gly Ala Trp Cys Ala Glu Glu Gln Asp Ala Asp Pro Trp Phe Gln Val
118 165 170 175

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120 Asp Ala Gly His Pro Thr Arg Phe Ser Gly Val Ile Thr Gln Gly Arg
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123 Asn Ser Val Trp Arg Tyr Asp Trp Val Thr Ser Tyr Lys Val Gln Phe
124          195          200          205
126 Ser Asn Asp Ser Arg Thr Trp Gly Ser Arg Asn His Ser Ser Gly
127          210          215          220
129 Met Asp Ala Val Phe Pro Ala Asn Ser Asp Pro Glu Thr Pro Val Leu
130 225          230          235          240
132 Asn Leu Leu Pro Glu Pro Gln Val Ala Arg Phe Ile Arg Leu Leu Pro
133          245          250          255
135 Gln Thr Trp Leu Gln Gly Gly Ala Pro Cys Leu Arg Ala Glu Ile Leu
136          260          265          270
138 Ala Cys Pro Val Ser Asp Pro Asn Asp Leu Phe Leu Glu Ala Pro Ala
139          275          280          285
141 Ser Gly Ser Ser Asp Pro Leu Asp Phe Gln His His Asn Tyr Lys Ala
142          290          295          300
144 Met Arg Lys Leu Met Lys Gln Val Gln Glu Gln Cys Pro Asn Ile Thr
145 305          310          315          320
147 Arg Ile Tyr Ser Ile Gly Lys Ser Tyr Gln Gly Leu Lys Leu Tyr Val
148          325          330          335
150 Met Glu Met Ser Asp Lys Pro Gly Glu His Glu Leu Gly Glu Pro Glu
151          340          345          350
153 Val Arg Tyr Val Ala Gly Met His Gly Asn Glu Ala Leu Gly Arg Glu
154          355          360          365
156 Leu Leu Leu Leu Leu Met Gln Phe Leu Cys His Glu Phe Leu Arg Gly
157          370          375          380
159 Asn Pro Arg Val Thr Arg Leu Leu Ser Glu Met Arg Ile His Leu Leu
160 385          390          395          400
162 Pro Ser Met Asn Pro Asp Gly Tyr Glu Ile Ala Tyr His Arg Gly Ser
163          405          410          415
165 Glu Leu Val Gly Trp Ala Glu Gly Arg Trp Asn Asn Gln Ser Ile Asp
166          420          425          430
168 Leu Asn His Asn Phe Ala Asp Leu Asn Thr Pro Leu Trp Glu Ala Gln
169          435          440          445
171 Asp Asp Gly Lys Val Pro His Ile Val Pro Asn His His Leu Pro Leu
172          450          455          460
174 Pro Thr Tyr Tyr Thr Leu Pro Asn Ala Thr Val Ala Pro Glu Thr Arg
175 465          470          475          480
177 Ala Val Ile Lys Trp Met Lys Arg Ile Pro Phe Val Leu Ser Ala Asn
178          485          490          495
180 Leu His Gly Gly Glu Leu Val Val Ser Tyr Pro Phe Asp Met Thr Arg
181          500          505          510
183 Thr Pro Trp Ala Ala Arg Glu Leu Thr Pro Thr Pro Asp Asp Ala Val
184          515          520          525
186 Phe Arg Trp Leu Ser Thr Val Tyr Ala Gly Ser Asn Leu Ala Met Gln
187          530          535          540
189 Asp Thr Ser Arg Arg Pro Cys His Ser Gln Asp Phe Ser Val His Gly
190 545          550          555          560
192 Asn Ile Ile Asn Gly Ala Asp Trp His Thr Val Pro Gly Ser Met Asn

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193          565          570          575
195 Asp Phe Ser Tyr Leu His Thr Asn Cys Phe Glu Val Thr Val Glu Leu
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198 Ser Cys Asp Lys Phe Pro His Glu Asn Glu Leu Pro Gln Glu Trp Glu
199          595          600          605
201 Asn Asn Lys Asp Ala Leu Leu Thr Tyr Leu Glu Gln Val Arg Met Gly
202          610          615          620
204 Ile Ala Gly Val Val Arg Asp Lys Asp Thr Glu Leu Gly Ile Ala Asp
205 625          630          635          640
207 Ala Val Ile Ala Val Asp Gly Ile Asn His Asp Val Thr Thr Ala Trp
208          645          650          655
210 Gly Gly Asp Tyr Trp Arg Leu Leu Thr Pro Gly Asp Tyr Met Val Thr
211          660          665          670
213 Ala Ser Ala Glu Gly Tyr His Ser Val Thr Arg Asn Cys Arg Val Thr
214          675          680          685
216 Phe Glu Glu Gly Pro Phe Pro Cys Asn Phe Val Leu Thr Lys Thr Pro
217          690          695          700
219 Lys Gln Arg Leu Arg Glu Leu Leu Ala Ala Gly Ala Lys Val Pro Pro
220 705          710          715          720
222 Asp Leu Arg Arg Arg Leu Glu Arg Leu Arg Gly Gln Lys Asp
223          725          730

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226 <210> SEQ ID NO: 3

227 <211> LENGTH: 20190

228 <212> TYPE: DNA

229 <213> ORGANISM: Homo sapiens

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234 <223> OTHER INFORMATION: Wherein n is A, C, G, or T

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238 <222> LOCATION: (2207)

239 <223> OTHER INFORMATION: Wherein n is A, C, G, or T

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243 <222> LOCATION: (2209)

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248 <222> LOCATION: (2211)

249 <223> OTHER INFORMATION: Wherein n is A, C, G, or T

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254 <223> OTHER INFORMATION: Wherein n is A, C, G, or T

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258 <222> LOCATION: (2215)

259 <223> OTHER INFORMATION: Wherein n is A, C, G, or T

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Input Set : A:\Cg-81cil.app

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272 <221> NAME/KEY: misc_feature
273 <222> LOCATION: (2221)
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277 <221> NAME/KEY: misc_feature
278 <222> LOCATION: (2222)
279 <223> OTHER INFORMATION: Wherein n is A, C, G, or T
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284 <223> OTHER INFORMATION: Wherein n is A, C, G, or T
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287 <221> NAME/KEY: misc_feature
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289 <223> OTHER INFORMATION: Wherein n is A, C, G, or T
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292 <221> NAME/KEY: misc_feature
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294 <223> OTHER INFORMATION: Wherein n is A, C, G, or T
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297 <221> NAME/KEY: misc_feature
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299 <223> OTHER INFORMATION: Wherein n is A, C, G, or T
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302 <221> NAME/KEY: misc_feature
303 <222> LOCATION: (2230)
304 <223> OTHER INFORMATION: Wherein n is A, C, G, or T
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309 accccggccc tgcatagcag cccggcacag ccgcccggcg agacagctaa cgggacctca 180
310 gaacagcatg tccggattcg agtcatcaag aagaaaaagg tcattatgaa gaagcggaag 240
311 aagctaactc taactcgcgc caccctactg gtgactgccg ggccccttgt gacccctact 300
312 ccagcaggga cctcgcaccc cgctgagaaa caagaaacag gctgtcctcc tttgggtctg 360
313 gagtccctgc gagtttcaga tagccggctt gaggcatcca gcagccagtc ctttgggtctt 420
314 ggaccacacc gaggacggct caacattcag tcaggcctgg aggacggcga tctatatgat 480
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317 gtcacatcat acaaggtcca gttcagcaat gacagtcgga cctgggtggg aagtaggaac 660
318 cacagcagtg ggatggacgc agtatttctt gccaatcag acccagaaac tccagtgtct 720
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RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:3; N Pos. 2227,2229,2230